

# Group B *Streptococcus* and *Streptococcus suis* Capsular Polysaccharides Induce Chemokine Production by Dendritic Cells via Toll-Like Receptor 2- and MyD88-Dependent and -Independent Pathways

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*Streptococcus agalactiae* (also known as group B *Streptococcus* [GBS]) and *Streptococcus suis* are encapsulated streptococci causing severe septicemia and meningitis. Bacterial capsular polysaccharides (CPSs) are poorly immunogenic, but anti-CPS antibodies are essential to the host defense against encapsulated bacteria. The mechanisms underlying anti-CPS antibody responses are not fully elucidated, but the biochemistry of CPSs, particularly the presence of sialic acid, may have an immunosuppressive effect. We investigated the ability of highly purified *S. suis* and GBS native (sialylated) CPSs to activate dendritic cells (DCs), which are crucial actors in the initiation of humoral immunity. The influence of CPS biochemistry was studied using CPSs extracted from different serotypes within these two streptococcal species, as well as desialylated CPSs. No interleukin-1 $\beta$  (IL-1 $\beta$ ), IL-6, IL-12p70, tumor necrosis factor alpha (TNF- $\alpha$ ), or IL-10 production was observed in *S. suis* or GBS CPS-stimulated DCs. Moreover, these CPSs exerted immunosuppressive effects on DC activation, as a diminution of gamma interferon (IFN- $\gamma$ )-induced B cell-activating factor of the tumor necrosis factor family (BAFF) expression was observed in CPS-pretreated cells. However, *S. suis* and GBS CPSs induced significant production of CCL3, via partially Toll-like receptor 2 (TLR2)- and myeloid differentiation factor 88 (MyD88)-dependent pathways, and CCL2, via TLR-independent mechanisms. No major influence of CPS biochemistry was observed on the capacity to induce chemokine production by DCs, indicating that DCs respond to these CPSs in a patterned way rather than a structure-dedicated manner.

*Streptococcus agalactiae* (also known as group B *Streptococcus* [GBS]) is a major cause of life-threatening invasive bacterial infections in pregnant women and neonates as well as in the elderly and immunocompromised individuals (1, 2). Clinical manifestations are mainly pneumonia, septicemia, and meningitis. Among 10 GBS serotypes identified, type III is one of the major serotypes associated with invasive neonatal infection and is the most common type in GBS meningitis (2). In addition, GBS type V is emerging as a leading cause of invasive disease in adults (3). *Streptococcus suis* is an important swine pathogen and an emerging zoonotic pathogen in humans that is able to induce septicemia with sudden death, meningitis, endocarditis, pneumonia, and arthritis (4, 5). Of the 35 serotypes, type 2 is the most virulent and is frequently isolated from both swine and humans (6), and type 14 is also emerging as a zoonotic threat (7). For both pathogens, the capsular polysaccharide (CPS), which defines the serotype, is considered the major virulence factor (8, 9). The structures of type III and V GBS CPSs are formed by different arrangements of the monosaccharides glucose, galactose, and *N*-acetylglucosamine into unique repeating units that contain a side chain terminated by sialic acid (*N*-acetylneuraminic acid [Neu5Ac]). The structures of type 2 and 14 *S. suis* CPSs are composed of the monosaccharides glucose, galactose, *N*-acetylglucosamine, and rhamnose (for type 2 only) arranged into a unique repeating unit that also contains a side chain terminated by sialic acid. In fact, these streptococci are the sole Gram-positive bacteria possessing sialic acid in their capsules (8, 10, 11). However, despite similarities in the compositions of the CPSs of these two bacterial species, each CPS is composed of a unique arrangement of these sugars conferring a distinct antigenicity. Moreover, sialic acid forms an  $\alpha$ -2,6 linkage with the adja-

cent galactose in *S. suis*, in contrast to the  $\alpha$ -2,3 linkage in GBS. Interestingly, the interplay of CPSs with components of the immune system, including antigen-presenting cells (APCs), seems to differ radically. Experiments using nonencapsulated mutants have shown that *S. suis* type 2 CPS has a strong antiphagocytic effect, in contrast to GBS type III CPS, and severely interferes with the release of most of the cytokines produced by *S. suis*-infected APCs. In the case of GBS type III, cytokine production is only partially modified or unaltered by the presence of its CPS (12–15).

Sialic acid of bacterial polysaccharides has been suggested to be involved in immune evasion via several mechanisms. For example, sialic acid of GBS type III CPS interferes with the immune response by molecular mimicry (8) and inhibition of complement activation (16). Some receptors expressed on the surfaces of leukocytes have a distinct preference for specific types of linkage of sialic acid to subterminal sugars. As these binding preferences are likely related to their biological functions, differences in sialic acid linkage in *S. suis* versus GBS might differentially modulate host

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immune responses (17). However, knowledge on the specific contributions of sialic acid to the interactions of these two pathogens with the immune system is restricted by the fact that deletion of genes involved in sialic acid synthesis results in considerable or complete loss of CPS expression at the bacterial surface (18, 19).

Dendritic cells (DCs), the most powerful APCs, express a wide variety of pattern recognition receptors (PRRs) that enable them to detect the presence of several pathogens through the recognition of pathogen-associated molecular patterns. Among these PRRs, Toll-like receptors (TLRs) are important for the initiation of the immune response as well as the shaping of adaptive immunity (20). The interactions between DCs and pathogens can strongly influence the magnitude and phenotype of the ensuing cellular and humoral adaptive immune responses, notably via the release of cytokines (21). Purified bacterial CPSs are classically reported to be T cell-independent (TI) antigens which are consequently weak stimulators of the host immune response (22, 23). However, several *in vitro* studies have demonstrated the ability of bacterial CPSs to interact with APCs, resulting in the production of cytokines and chemokines (24–28). The adaptor molecule myeloid differentiation factor 88 (MyD88), which is involved in intracellular events downstream of TLR signaling, and TLR2 have been suspected to be involved in the interactions of bacterial CPSs with DCs and/or macrophages (25, 27). Nevertheless, the potential role of pure carbohydrates as ligands for TLRs, and more globally for PRRs, remains largely uninvestigated.

Recent reports have shown that DCs play an important role in TI responses and, more precisely, in the development of the humoral response, via the release of B cell-activating factor of the tumor necrosis factor family (BAFF). BAFF is able to enhance B cell proliferation, immunoglobulin (Ig) class switching, and Ig secretion (29, 30). These critical signals are particularly interesting in the context of infection by encapsulated bacteria, where antibodies (Abs) against the CPS have been proven to be essential to the host defense (23). As such, we hypothesized that intra- and interspecies structural differences in CPS might differently modulate DC release of cytokines and chemokines. The goal of this study was thus to evaluate and compare the effects of highly purified CPS preparations from *S. suis* types 2 and 14 as well as GBS types III and V on DC activation and, more specifically, on the capacity to induce cytokines essential for the development of an effective humoral immune response. The influence of sialic acid was analyzed using chemically desialylated CPS preparations.

## MATERIALS AND METHODS

**Native CPS purification.** The reference strains of *S. suis* serotype 2 (S735; ATCC 43765), isolated from a pig with meningitis (31), and *S. suis* serotype 14 (DAN13730), isolated from a human with meningitis (32), were grown in 150 ml of Todd-Hewitt broth (THB) (Oxoid, Thermo Fisher Scientific) at 37°C for 16 h, diluted to 6 liters in fresh THB, and grown to an optical density at 540 nm ( $OD_{540}$ ) of 0.8. The cells were pelleted by centrifugation at  $10,000 \times g$  for 40 min, suspended by repeated pipetting in 33 mM phosphate-buffered saline (PBS), pH 8.0, and chilled. The CPSs were then purified as previously described (10, 11).

GBS serotype III strain COH-1, isolated from an infant with bacteremia (33), and GBS serotype V strain CJB111 (ATCC BAA-23), isolated from a neonate with septicemia, were used in this study. GBS CPSs were prepared as previously reported (34), with some modifications. Briefly, bacteria were grown in 200 ml THB at 37°C for 16 h, diluted to 8 liters in fresh THB, and grown to an  $OD_{540}$  of 0.8. The cells were pelleted by centrifugation at  $10,000 \times g$  for 40 min, washed in PBS, pH 7.3, and

treated with 1 N NaOH at 37°C overnight. After neutralization and dialysis, proteins were digested by treatment with 1 mg/ml pronase (Sigma-Aldrich) at 37°C overnight, followed by dialysis. The CPSs were then subjected to re-*N*-acetylation with 0.8 M acetic anhydride (Sigma) in 5 N NaOH and finally purified by gel filtration on Sephacryl S-300 (GE Healthcare), using 50 mM  $NH_4HCO_3$  as the eluent.

**CPS desialylation.** Highly purified native CPSs were desialylated by mild acid hydrolysis as described previously (10). Briefly, CPS (8 mg) was heated in 1 ml of 70 mM HCl at 60°C for 250 min, neutralized with 2 M  $NH_4OH$ , dialyzed against deionized water for 48 h at 4°C with a Spectra/Por membrane (molecular size cutoff of 3,500 Da; Spectrum Laboratories), and freeze-dried.

**CPS quality controls.** Each purified CPS was subjected to rigorous quality control tests as previously described (10). Nucleic acids were quantified using an ND 1000 spectrometer (Nanodrop). The absorbance was measured at 230 and 260 nm. Calculations were done with Nanodrop software. According to the manufacturer, results are reproducible between 2 and 100 ng/ $\mu$ l. Proteins were quantified by use of a modified Lowry protein assay kit from Pierce on 1-mg/ml CPS samples, using a standard curve prepared with diluted albumin standards from 1 to 1,000  $\mu$ g/ml. The calculated limit of detection ( $P \leq 0.05$ ) was 0.7 to 1.3  $\mu$ g/ml. Each CPS was analyzed by nuclear magnetic resonance (NMR) as described below. The monosaccharide composition of polysaccharides was confirmed by methanolysis followed by acetylation and analysis by gas chromatography (GC), either with flame ionization detection or coupled to mass spectrometry as previously described (10). The weight-average molecular weight ( $M_w$ ) of each CPS was determined by size-exclusion chromatography coupled with multiangle light scattering (SEC-MALS) as described below. The presence (native CPS) or absence (desialylated CPS) of sialic acid was verified by NMR and by an enzyme-linked lectin assay (ELLA) as described below.

**Nuclear magnetic resonance assay.** *S. suis* native CPSs were exchanged in phosphate buffer ( $p^2H$  8.0) in  $^2H_2O$  (99.9 atom%  $^2H$ ), freeze-dried, and dissolved in  $^2H_2O$  (99.96 atom%  $^2H$ ) to a final phosphate concentration of 33 mM. The other CPSs were exchanged in  $^2H_2O$  (99.9 atom%  $^2H$ ), freeze-dried, and dissolved in  $^2H_2O$  (99.96 atom%  $^2H$ ). NMR spectra were acquired on CPS samples at concentrations of ca. 1% to 2%. Conventional  $^1H$  spectra were acquired at 14 T on Bruker Avance spectrometers equipped with either a 5-mm TCI CryoProbe at 50°C or a 5-mm PABBO BB inverse gradient probe at 75°C or at 11.75 T on a Bruker Avance 500 spectrometer equipped with a 5-mm triple-resonance TBI probe at 60 to 80°C, using standard Bruker pulse sequences.

**$M_w$  characterization of native and desialylated CPSs.** The  $M_w$  of CPSs were characterized by SEC-MALS. Chromatographic separation was performed with two 8-mm by 300-mm Shodex OHPak gel filtration columns connected in series (SB 806 and SB 804), preceded by an SB 807G guard column (Showa Denko). Elution was done with a Waters 510 pump (Waters), using a 0.1 M  $NaNO_3$  mobile phase filtered through a 0.02- $\mu$ m membrane (Whatman), at a flow rate of 0.5 ml/min. Samples were dissolved in the SEC eluent at concentrations of 0.7 to 1.0 mg/ml for native CPSs and 2.0 to 3.5 mg/ml for desialylated CPSs and then were injected with a 100- or 200- $\mu$ l sample loop. Molecular masses were determined with a Dawn EOS MALS detector (Wyatt). A model RI 410 differential refractometer (Waters) was used as a concentration detector. A refractive index increment ( $dn/dc$ ) of 0.137 ml/g was calculated for 690 nm, using data for xanthan at 436 and 546 nm (35), and the second virial coefficient ( $A_2$ ) was taken as zero. Calculations were performed with ASTRA software, version 6.0.0.108 (Wyatt).

**ELLA.** In order to verify the presence or absence of sialic acid in the purified native and desialylated CPSs, ELLA was carried out based on a previously described technique (36), which was adapted to CPSs. Briefly, 200 ng of sample (native or desialylated CPS) was added to wells of an enzyme-linked immunosorbent assay (ELISA) plate (Nunc-Immuno Polysorp). After overnight coating at 4°C, the wells were washed and blocked by the addition of  $1 \times$  Carbo-Free solution (Vector Laboratories).

After washings, the wells were incubated for 1 h with biotinylated *Sam-bucus nigra* agglutinin (SNA-I) (Vector Laboratories), which specifically recognizes sialic acid as Neu5Ac $\alpha$ -2,6-Galp/GalpNAc (37), or biotinylated *Maackia amurensis* leucoagglutinin (MAL-I) (Vector Laboratories), which recognizes sialic acid as Neu5Ac $\alpha$ -2,3-Gal $\beta$ -1,4-GlcNAc (38). Horseradish peroxidase (HRP)-labeled avidin D (Vector Laboratories) and 3,3',5,5'-tetramethylbenzidine were then added. In some experiments, HRP-conjugated *Limax flavus* agglutinin (LFA) (Vector Laboratories), which recognizes Neu5Ac (39), was used. The enzyme reaction was stopped by the addition of 0.5 M H<sub>2</sub>SO<sub>4</sub>, and the absorbance was read at 450 nm with an ELISA plate reader.

**Dot-ELISA.** Ten microliters of purified native or desialylated CPS (each at 1 mg/ml) or 10  $\mu$ l of heat-killed whole bacteria was blotted on a polyvinylidene difluoride (PVDF) Western blot membrane (Roche). Heat-killed bacteria were obtained after incubating bacteria at 60°C for 45 min and were adjusted to 10<sup>9</sup> CFU/ml. The membrane was blocked for 1 h with a solution of Tris-buffered saline (TBS) containing 2% casein, followed by 2 h of incubation with either the mouse monoclonal Ab (MAb) Z3, which specifically recognizes the sialic acid moiety of *S. suis* type 2 CPS (40); monospecific polyclonal rabbit sera against *S. suis* type 2 (41) or *S. suis* type 14 (32); or commercial rabbit sera against GBS type III or GBS type V CPS (Denka Seiken). The membrane was washed, and the appropriate anti-rabbit or anti-mouse HRP-conjugated Ab (Jackson) was added for 1 h. The membrane was washed 3 times with TBS and revealed with a 4-chloro-1-naphthol solution (Sigma).

**Mouse strains and generation of bone marrow-derived dendritic cells.** Six- to 8-week-old mice originating from Jackson Laboratory, including wild-type (WT) C57BL/6, MyD88<sup>-/-</sup> (B6.129P2-Myd88<sup>tm1Defr/J</sup>), and TLR2<sup>-/-</sup> (B6.129-Tlr2<sup>tm1Kir/J</sup>) mice, were used. All experiments involving mice were conducted in accordance with the guidelines and policies of the Canadian Council on Animal Care and the principles set forth in the *Guide for the Care and Use of Laboratory Animals* by the Animal Welfare Committee of the Université de Montréal (42). Bone marrow-derived DCs were produced according to a previously described technique (13, 43) and cultured in complete medium consisting of RPMI 1640 supplemented with 5% heat-inactivated fetal bovine serum, 10 mM HEPES, 2 mM L-glutamine, 50  $\mu$ M 2-mercaptoethanol, 100 U/ml penicillin-streptomycin, and 20  $\mu$ g/ml gentamicin. All reagents were from Gibco (Invitrogen). Cell purity was routinely  $\geq$ 86 to 90% CD11c<sup>high</sup> F4/80<sup>-dim</sup> cells as determined by fluorescence-activated cell sorter (FACS) analysis, in agreement with values reported in other studies (44–46).

**In vitro DC stimulation assay.** DCs were resuspended at 10<sup>6</sup> cells/ml in complete medium and stimulated with native or desialylated CPS (5, 50, 100, or 200  $\mu$ g/ml). At 6 and 24 h, supernatants were collected for cytokine quantification by ELISA, and cells were harvested for analysis of BAFF expression by reverse transcriptase quantitative PCR (RT-qPCR). Cells stimulated with 1  $\mu$ g/ml ultrapurified *Escherichia coli* O55:B5 lipopolysaccharide (LPS) (Apotech Corporation) and 10 ng/ml recombinant mouse gamma interferon (IFN- $\gamma$ ) (R&D Systems) served as positive controls for cytokine production and BAFF expression, respectively. Non-stimulated cells served as a negative control. In some experiments, DCs were prestimulated with CPS (at 200  $\mu$ g/ml) for 6 h prior to incubation with 10 ng/ml IFN- $\gamma$  for 24 h. Cells were then harvested for BAFF expression analysis. DCs preincubated in complete medium before addition of IFN- $\gamma$  served as a control. All solutions and CPSs were tested for the absence of endotoxin by use of a *Limulus* amoebocyte lysate gel-clotting test (Pyrotell) with a sensitivity limit of 0.03 endotoxin unit (EU)/ml. Otherwise, the absence of endotoxin contamination during cell stimulation was controlled by parallel assays with polymyxin B sulfate (Sigma) at 20  $\mu$ g/ml.

**Cytokine quantification by ELISA.** Levels of interleukin-1 $\beta$  (IL-1 $\beta$ ), IL-6, IL-10, IL-12p70, tumor necrosis factor alpha (TNF- $\alpha$ ), CCL2 (MCP-1), and CCL3 (MIP-1 $\alpha$ ) in cell culture supernatants were measured by sandwich ELISAs using pair-matched Abs from R&D Systems,

TABLE 1 Quality control tests of purified *S. suis* and GBS CPSs<sup>f</sup>

CPS <sup>a</sup>	Sugar composition <sup>b</sup>	M <sub>w</sub> <sup>c</sup>	% (wt/wt) nucleic acids <sup>d</sup>	% (wt/wt) protein <sup>e</sup>
<i>S. suis</i> type 2 (n)	1:2.6:0.9:1.0:0.9	480	0.8	NS
<i>S. suis</i> type 2 (dS)	1:2.6:0.9:1.0:0.9	21	0.8	NS
<i>S. suis</i> type 14 (n)	1:2.5:0.8:1.0:0.0	500	0.7	NS
<i>S. suis</i> type 14 (dS)	1:2.5:0.8:1.0:0.0	176	0.7	NS
GBS type III (n)	1:1.5:1.0:0.9:0.0	108	0.8	NS
GBS type III (dS)	1:1.5:1.0:0.9:0.0	59	0.8	NS
GBS type V (n)	3:1.5:0.8:0.8:0.0	128	0.9	NS
GBS type V (dS)	3:1.5:0.8:0.8:0.0	91	0.9	NS

<sup>a</sup> n, native; dS, desialylated.

<sup>b</sup> Determined by GC after methanolysis and acetylation and shown in the order glucose: galactose:N-acetylglucosamine:N-acetylneuraminic acid:rhamnose.

<sup>c</sup> Determined by SEC-MALS (expressed in kg/mol).

<sup>d</sup> Determined by spectrophotometry at 230 and 260 nm.

<sup>e</sup> Determined by spectrophotometry at 750 nm. NS, nonsignificant (under the limit of detection).

<sup>f</sup> See Materials and Methods for more details.

performed according to the manufacturer's recommendations. Twofold dilutions of recombinant mouse cytokines were used to generate standard curves. Sample dilutions giving OD readings in the linear portion of the appropriate standard curve were used to quantify the levels of each cytokine.

**Analysis of BAFF gene expression by real-time RT-qPCR.** Total RNA was isolated from 10<sup>6</sup> DCs by use of TRIzol reagent (Invitrogen). After elimination of genomic DNA, 800 ng of total RNA was reverse transcribed with a QuantiTect reverse transcription kit (Qiagen). The cDNA was amplified using an SsoFast EvaGreen Supermix kit (Bio-Rad). The PCR amplification program for all cDNAs consisted of an enzyme activation step of 3 min at 98°C followed by 40 cycles of a denaturing step of 2 s at 98°C and an annealing/extension step of 5 s at 56°C. The ATP synthase subunit beta (ATP5B) and cytochrome c<sub>1</sub> (CYC1) genes were used as normalizing genes to compensate for potential differences in amounts of cDNA. The primers used for amplification of the different target cDNAs are listed in Table S1 in the supplemental material and were all tested to achieve amplification efficiencies between 98.5% and 100%. The primer sequences were all designed from the NCBI GenBank mRNA sequence, using the Web-based software primerquest from Integrated DNA Technologies. A Bio-Rad CFX-96 sequence detector was used for amplification of cDNA, and quantitation of differences between the different groups was calculated using the 2<sup>- $\Delta\Delta$ CT</sup> method (47). Nonstimulated DCs were used as the calibrator reference in the analysis.

**Statistical analysis.** All data are expressed as means  $\pm$  standard errors of the means (SEM). Data were analyzed for significance using analysis of variance (ANOVA). Significance is denoted in the figures as follows: \*,  $P < 0.05$ ; and \*\*,  $P < 0.001$ . All experiments were repeated at least three times.

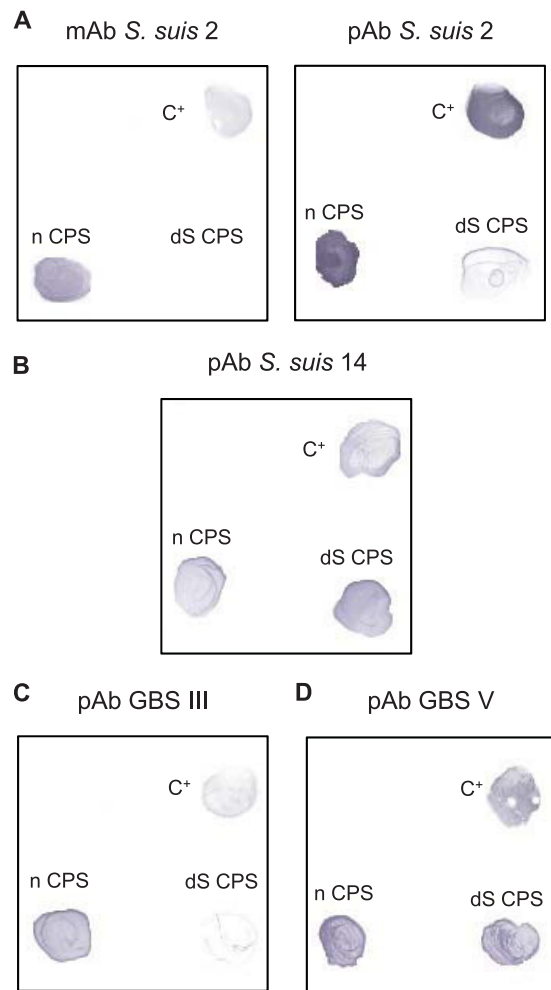
## RESULTS

***S. suis* and GBS CPS purification, desialylation, and quality control tests.** On average, from a 6-liter *S. suis* type 2 or *S. suis* type 14 culture, 150 mg of crude capsule was obtained, to afford around 30 mg of purified CPS after gel filtration (average yield of 5 mg CPS/liter of culture). From 8 liters of GBS type III or type V culture, 360 mg or 600 mg initial crude capsule was obtained, respectively, to afford around 30 mg of purified CPS after gel filtration (average yield of 3.75 mg CPS/liter of culture). GC and NMR analyses of purified CPSs gave sugar compositions and structures in accord with previous findings (8, 10, 11) (Table 1; see Fig. S1 and S2 in the supplemental material). Each repeating unit is com-

posed of the same four sugars (with an additional rhamnose in *S. suis* type 2 CPS), with sialic acid located at the terminal side chain. The ratios of the four common sugars are similar in *S. suis* type 2 and 14 CPSs, whereas they are different between GBS type III and V CPSs. In comparison with *S. suis* CPSs, the repeating unit of GBS type III and V CPSs has one less galactose, and GBS type V CPS has two additional glucose residues. Indeed, the glucose:galactose:*N*-acetylglucosamine:*N*-acetylneuraminic acid:rhamnose sugar ratios for each repeating unit of *S. suis* type 2, *S. suis* type 14, GBS type III, and GBS type V CPSs are 1:3:1:1:1, 1:3:1:1:0, 1:2:1:1:0, and 3:2:1:1:0, respectively (Table 1; see Fig. S1 and S2). One of the problems encountered in purifying GBS CPSs is contamination with group B antigen, as the CPS is covalently linked to the cell wall in this bacterial species (48). The absence of rhamnose (a sugar present in group B antigen but not in GBS CPSs) in purified GBS CPS preparations confirmed the absence of contamination with this cell wall antigen. No protein was found above the limit of detection, indicating that there was less than 1.3% (wt/wt) protein in all purified CPSs, and DNA and RNA contamination was less than 1% (wt/wt) (Table 1). SEC-MALS showed that native *S. suis* type 2 and 14 CPSs had similar  $M_w$ , which were 480 and 500 kg/mol, respectively. Native GBS type III and V CPSs had comparable  $M_w$ , which were almost 4-fold lower than that of native *S. suis* CPSs. Desialylation induced a larger  $M_w$  decrease in *S. suis* CPSs than in GBS CPSs, with the highest diminution for desialylated *S. suis* type 2 CPS (96% of the initial  $M_w$ ) (Table 1). This could be explained by the presence of rhamnose in the backbone of *S. suis* type 2 CPS, which forms a linkage more susceptible to acid hydrolysis with the adjacent sugar.

***S. suis* and GBS CPS recognition by specific sera and sialic acid-binding lectins.** Dot-ELISA experiments on native CPSs showed that recognition of the CPS epitope of each CPS preparation was conserved after purification (Fig. 1). Whereas the native *S. suis* type 2 CPS was well recognized by MAb Z3, which is specific for the sialic acid part of the capsule, the negative reaction of the desialylated *S. suis* type 2 CPS attested the absence of sialic acid in the latter (Fig. 1A, left panel). Desialylation of *S. suis* type 2 CPS resulted in reduced recognition by specific polyclonal Abs (Fig. 1A, right panel), whereas recognition of desialylated preparations of *S. suis* type 14 CPS was unaltered (Fig. 1B). The capacity of specific polyclonal Abs to react with the desialylated GBS type III CPS was almost completely lost compared to that with native CPS (Fig. 1C). In contrast, native and desialylated preparations of GBS type V CPS were similarly recognized by specific polyclonal Abs (Fig. 1D). These data suggest that there are intra- and interspecies variations in the immunogenic properties exerted by the sialic acid moiety (49–51).

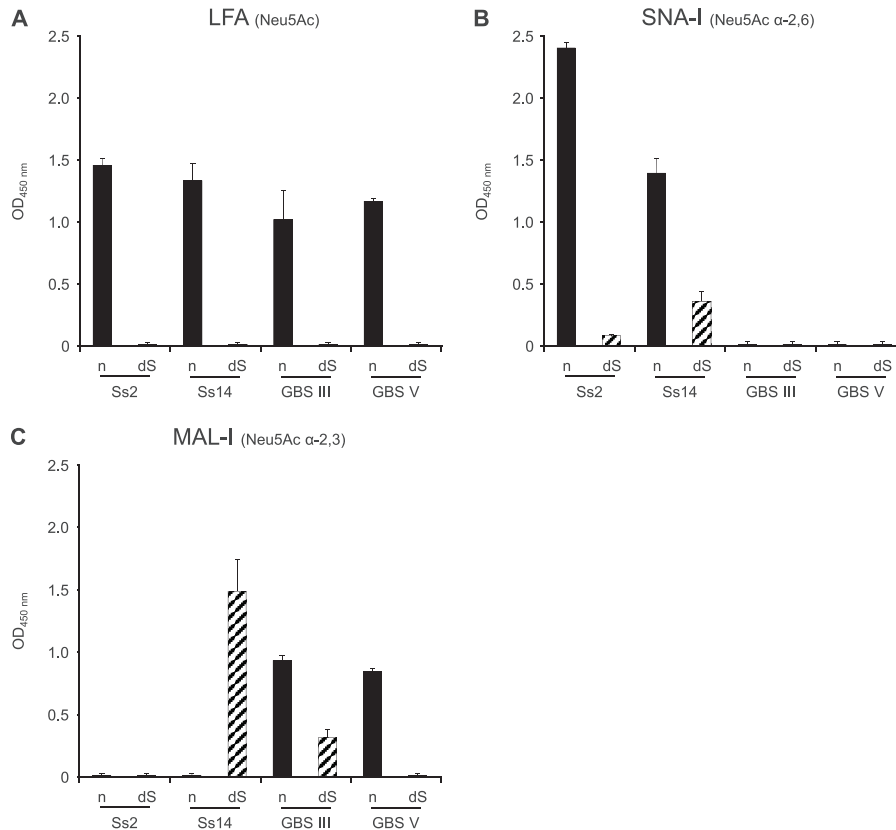
To further confirm the presence or absence of sialic acid in native and desialylated CPSs, an ELLA test was performed. Recognition of native *S. suis* and GBS CPSs by LFA, which is specific for Neu5Ac, confirmed the integrity of sialic acid in these preparations, whereas the absence of reaction with desialylated preparations demonstrated the absence of this sugar after desialylation by mild acid hydrolysis (Fig. 2A). Recognition of native *S. suis* CPSs by SNA-I and of native GBS CPSs by MAL-I validated that sialic acid forms  $\alpha$ -2,6 and  $\alpha$ -2,3 links, respectively, with the adjacent galactose (Fig. 2B and C). In the case of desialylated *S. suis* type 14 CPS, a positive reaction was observed with both SNA-I and MAL-I. Similarly, a positive reaction was observed for desialylated GBS type III CPS with MAL-I (Fig. 2B and C). Because negative



**FIG 1** Recognition of *S. suis* and GBS CPSs by dot-ELISA. *S. suis* type 2 (A), *S. suis* type 14 (B), GBS type III (C), or GBS type V (D) whole bacteria ( $10^7$  CFU) or their respective purified native (n) or desialylated (dS) CPSs (10  $\mu$ g) were incubated with a MAb directed against the sialic acid moiety of *S. suis* type 2 CPS (A, left) or with monospecific polyclonal sera (pAb) against *S. suis* type 2 CPS (A, right), *S. suis* type 14 CPS (B), GBS type III CPS (C), or GBS type V CPS (D). C<sup>+</sup>, positive control.

reactions with LFA and NMR analysis (Fig. 2A; see Fig. S1 and S2 in the supplemental material) clearly demonstrated the absence of sialic acid in these desialylated preparations, the positive reaction with SNA-I can be explained by the nonspecific binding of the lectin to D-galactose (37). Recognition of the Gal $\beta$ -1,4-GlcNAc epitope of desialylated *S. suis* type 14 and GBS type III CPSs by MAL-I can explain the unspecific reaction of this lectin with these two preparations (52).

***S. suis* and GBS CPSs induce the release of chemokines by DCs.** The levels of several cytokines and chemokines in the supernatants of DCs incubated with *S. suis* type 2 or 14 or GBS type III or V CPS (each at 200  $\mu$ g/ml) were measured 24 h after stimulation. No significant difference in pro- or anti-inflammatory cytokine production was observed between DCs incubated with the different CPSs and those incubated with medium alone (Fig. 3). Similar results were obtained 6 h after stimulation (data not shown). The presence of sialic acid did not influence the release of



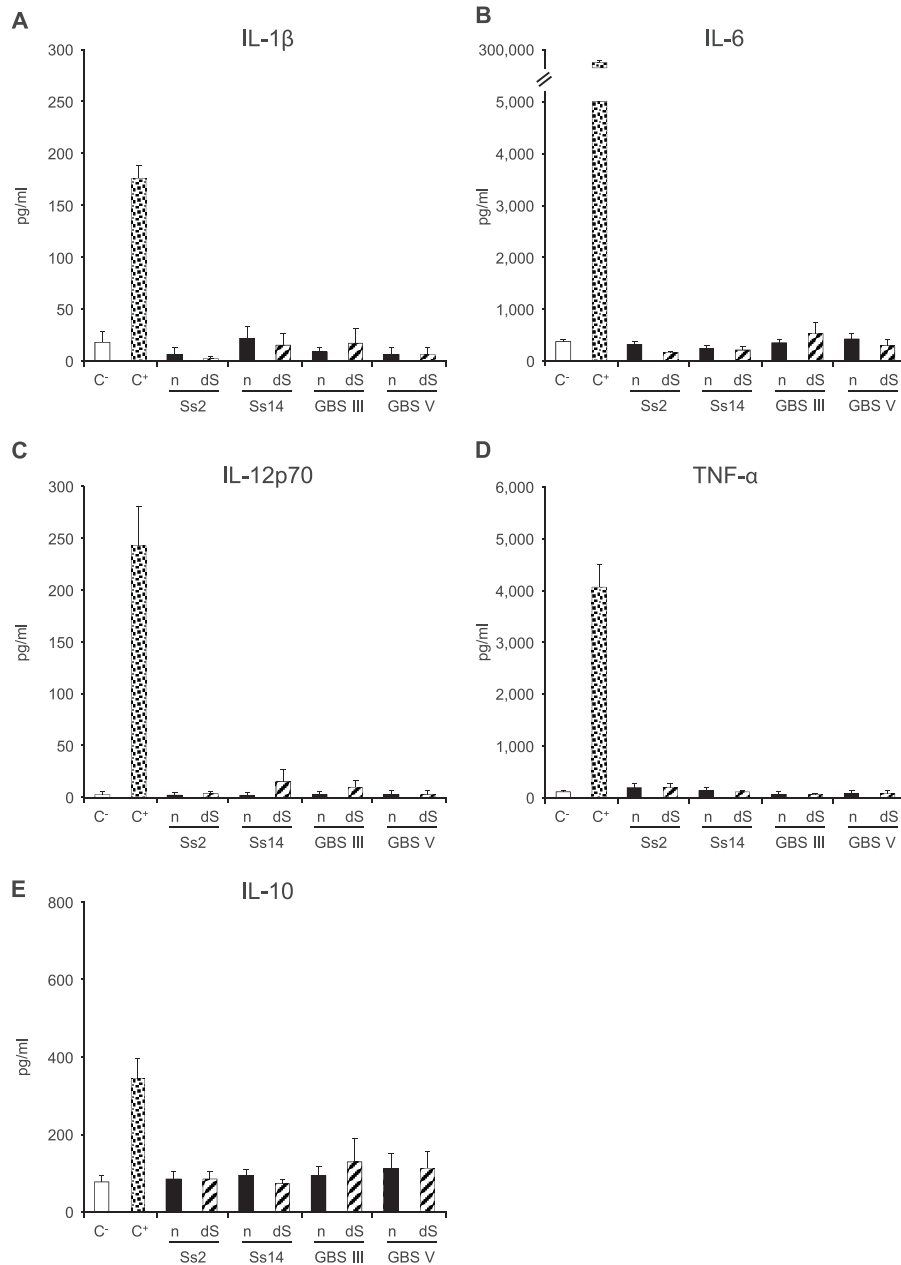
**FIG 2** Recognition of *S. suis* and GBS CPSs by ELLA. Native (n) or desialylated (dS) *S. suis* type 2 (Ss2), *S. suis* type 14 (Ss14), GBS type III, or GBS type V CPSs (each at 2  $\mu\text{g}/\text{ml}$ ) was incubated with *Limax flavus* agglutinin specific for Neu5Ac (LFA) (A), *Sambucus nigra* agglutinin specific for Neu5Ac  $\alpha$ -2,6 links (SNA-I) (B), or *Maackia amurensis* leucoagglutinin specific for Neu5Ac  $\alpha$ -2,3 links (MAL-I) (C). Data are expressed as mean OD<sub>450</sub> with SEM for at least three experiments with at least three technical replicates and are corrected for reaction of the dilution buffer with the corresponding lectin.

these cytokines, as no significant difference was observed between native and desialylated CPSs (Fig. 3).

In contrast, *S. suis* and GBS CPSs induced significant release of the chemokines CCL2 and CCL3 at 24 h (Fig. 4). CCL3 production was induced similarly by all CPS preparations (Fig. 4A). On the other hand, CCL2 production was significantly higher when DCs were activated with *S. suis* CPSs ( $P < 0.001$ ) (Fig. 4B). In the case of *S. suis*, more CCL2 production was observed for DCs stimulated with *S. suis* type 14 CPS than for those stimulated with *S. suis* type 2 CPS ( $P < 0.001$ ). Sialic acid plays a partial inhibitory role in the production of CCL2 for *S. suis* type 2 CPS only ( $P < 0.001$ ). To explore more precisely the characteristics of CCL2 and CCL3 production, dose- and time-response analyses were performed. CCL2 release induced by *S. suis* type 2 CPS was shown to be directly proportional to the CPS concentration and the time of incubation, with maximum release obtained at 24 h with a concentration of 200  $\mu\text{g}/\text{ml}$  CPS (Fig. 5). Similar results were obtained with *S. suis* type 14 and GBS type III and V CPSs, for both CCL2 and CCL3 production (data not shown).

**Involvement of PRRs in chemokine release by DCs stimulated with *S. suis* and GBS CPSs.** Previous *in vitro* studies have demonstrated that well-encapsulated *S. suis* type 2 induces TLR2 mRNA expression by human monocytes (53) and porcine DCs (15) and stimulates cytokine and chemokine production by murine macrophages and DCs in a TLR2- and MyD88-dependent manner (53, 54). Moreover, the release of cytokines by *S. suis*-

stimulated human monocytes was significantly reduced by Ab-mediated blocking of TLR2 but not TLR4 (53). In the case of GBS type III, the killed bacterium induces TLR2 mRNA expression and TLR2-dependent expression of certain cytokines and chemokines by murine macrophages (55). Killed GBS type III stimulates TNF- $\alpha$  release by those cells, in a TLR2-independent but MyD88-dependent way (56). As earlier studies have shown that PRRs such as TLRs are involved in cytokine production by APCs stimulated with purified CPSs from several different bacteria (25, 27, 57), we aimed to evaluate the implication of these receptors in the production of chemokines by DCs activated with our CPSs. Therefore, levels of CCL2 and CCL3 release were compared between WT and TLR2<sup>-/-</sup> or MyD88<sup>-/-</sup> DCs incubated with *S. suis* or GBS CPSs (at 200  $\mu\text{g}/\text{ml}$ ) for 24 h. As shown in Fig. 6A and 7A, no significant difference in CCL2 production was observed between WT and TLR2<sup>-/-</sup> DCs or between WT and MyD88<sup>-/-</sup> DCs for either *S. suis* or GBS CPSs. On the other hand, TLR2<sup>-/-</sup> DCs activated with either GBS type III or type V CPS showed a partial reduction of CCL3 production compared to their WT DC counterparts ( $P < 0.05$ ). In the case of *S. suis*, a significant effect of TLR2 on CCL3 production was observed only for native *S. suis* type 2 CPS, indicating the involvement of other receptors (Fig. 6B). Indeed, a significant impairment of CCL3 production by MyD88<sup>-/-</sup> DCs incubated with all four *S. suis* or GBS CPSs was observed, with decreases varying between 40 and 50% ( $P < 0.001$ ) (Fig. 7B). In general, the presence of sialic acid did not seem to

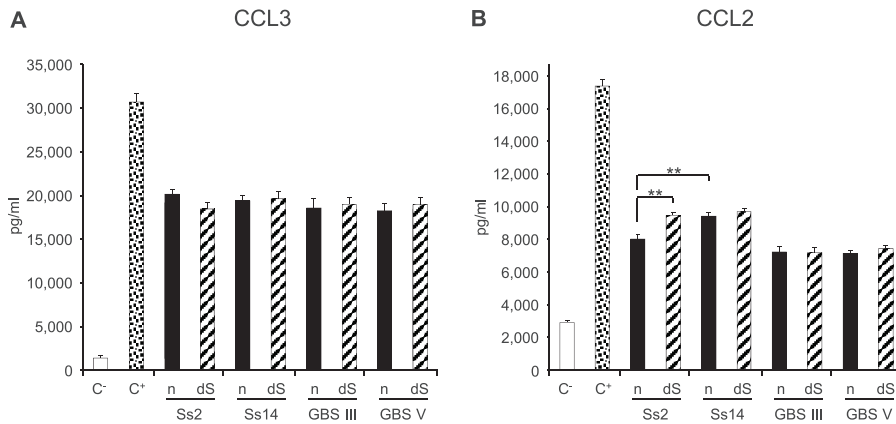


**FIG 3** Pro- and anti-inflammatory cytokine production by DCs in response to stimulation by *S. suis* or GBS CPSs for 24 h. Native (n) or desialylated (dS) *S. suis* type 2 (Ss2), *S. suis* type 14 (Ss14), GBS type III, or GBS type V CPS (each at 200  $\mu\text{g}/\text{ml}$ ) was incubated with DCs ( $10^6$  cells/ml). After 24 h, supernatants were collected, and IL-1 $\beta$  (A), IL-6 (B), IL-12p70 (C), TNF- $\alpha$  (D), and IL-10 (E) levels were quantified by ELISA. Cells stimulated with medium alone and with LPS (1  $\mu\text{g}/\text{ml}$ ) served as negative (C<sup>-</sup>) and positive (C<sup>+</sup>) controls, respectively. Data are expressed as means with SEM (pg/ml) for at least three experiments with at least three technical replicates.

significantly modulate the interactions between either GBS or *S. suis* CPSs and the TLR/MyD88 pathway (Fig. 6 and 7).

**Modulation of BAFF gene expression by DCs incubated with *S. suis* and GBS CPSs.** In contrast to the numerous studies that have focused on the effects of BAFF on B cell physiology, there is a relative paucity of evidence concerning BAFF production by DCs. A recent study has shown that the prototype TI antigen NP-Ficolin is able to induce the release of BAFF by murine DCs, which is essential for the development of Ab responses (58). To analyze whether *S. suis* and GBS CPSs modulate BAFF mRNA expression

levels, DCs were stimulated with native CPSs at 6 and 24 h. IFN- $\gamma$  (10 ng/ml), which is known to stimulate BAFF synthesis by monocytes, macrophages, and DCs (29, 59, 60), was used as a positive control. As shown in Fig. 8, no significant difference in BAFF expression levels was observed between CPS-activated DCs and control nonstimulated cells at either 6 or 24 h. Only a slight up-regulation (<2-fold increase) of BAFF mRNA expression was observed at 6 h for all native CPS preparations. As sialic acid has been shown to inhibit B cell activation (61), we wanted to know if this sugar could be involved in the inhibition of BAFF expression as



**FIG 4** Chemokine production by DCs in response to stimulation by *S. suis* or GBS CPSs for 24 h. Native (n) or desialylated (dS) *S. suis* type 2 (Ss2), *S. suis* type 14 (Ss14), GBS type III, or GBS type V CPS (each at 200  $\mu\text{g/ml}$ ) was incubated with DCs ( $10^6$  cells/ml). After 24 h, supernatants were collected, and CCL3 (A) and CCL2 (B) levels were quantified by ELISA. Cells stimulated with medium alone and with LPS (1  $\mu\text{g/ml}$ ) served as negative ( $C^-$ ) and positive ( $C^+$ ) controls, respectively. Data are expressed as means with SEM (pg/ml) for at least three experiments with at least three technical replicates. \*\*,  $P < 0.001$ .

well. Sialic acid did not significantly modulate BAFF expression by DCs (Fig. 8).

To further evaluate the capacity of GBS and *S. suis* CPSs to inhibit BAFF expression by DCs, we evaluated the impact of DC preincubation with CPS on IFN- $\gamma$ -induced BAFF. To this end, DCs were preincubated with the different CPSs for 6 h prior to stimulation with IFN- $\gamma$  for 24 h. Diminutions of BAFF mRNA expression ranging from 20 to 40% were observed when DCs were preincubated with either *S. suis* or GBS CPSs in comparison with nonpretreated cells, confirming the inhibitory effect of these bacterial CPSs on BAFF expression (Fig. 9).

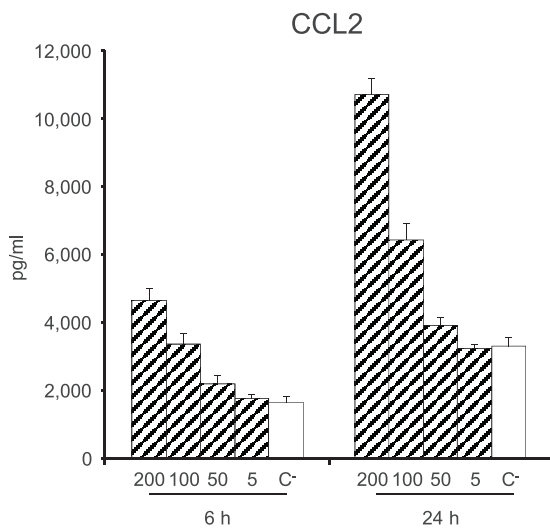
## DISCUSSION

CPS is a crucial component for both *S. suis* and GBS. In addition to forming the basis for serotype designation and being the major

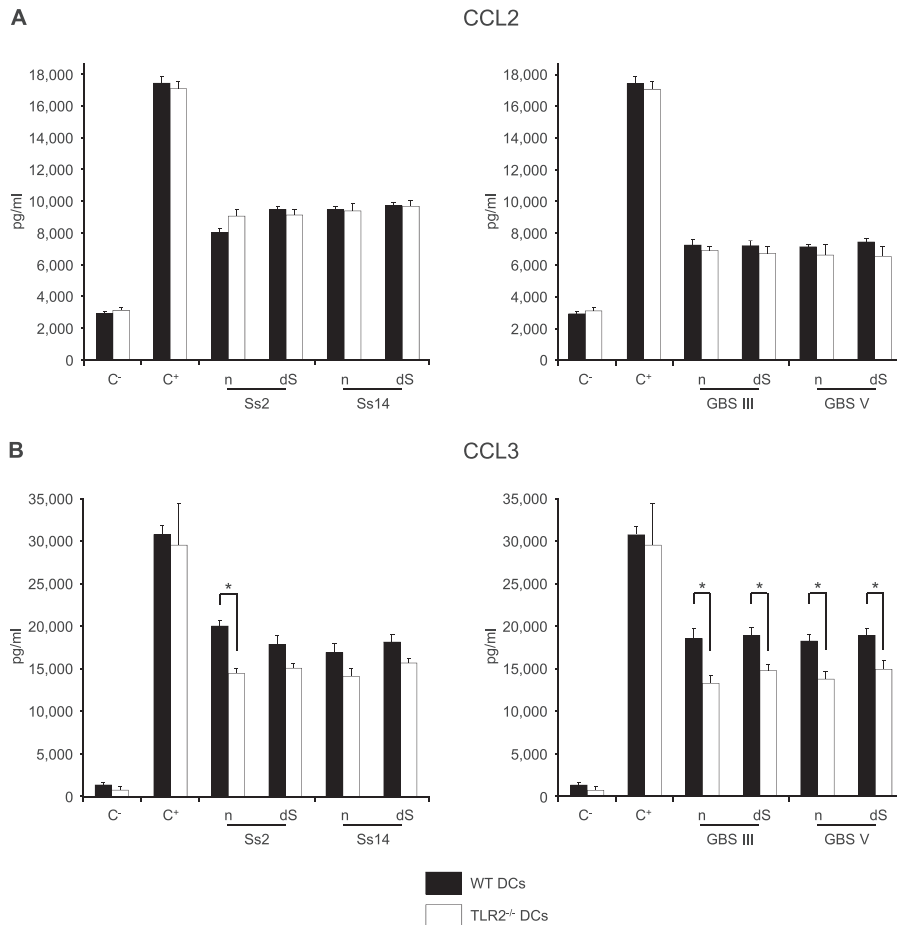
virulence factor, it has, as an immunogen, a high protective potential in the fight against infections by these two streptococci. Indeed, as shown with other encapsulated bacteria, such as *Streptococcus pneumoniae*, *Neisseria meningitidis*, and *Haemophilus influenzae*, the specific anti-CPS humoral response could play a decisive role in host survival of *S. suis* or GBS infections. Previous studies have demonstrated that mouse or pig anti-*S. suis* type 2 CPS Abs have a protective role in homologous opsonophagocytosis *in vitro* assays (40, 62, 63) or during *in vivo* challenge (40, 64). Finally, Abs raised against several GBS CPS serotypes, including serotypes Ia, Ib, II, III, and V, are protective against neonatal infections in both animals and humans and exhibit opsonophagocytosis ability *in vitro* (65).

However, *S. suis* and GBS CPSs are poorly immunogenic molecules because they cannot recruit T cell help for B cell functions. Mothers of neonates developing GBS type III disease have low concentrations of anti-CPS Abs in sera at delivery (66), and very few anti-CPS Abs can be detected in pigs infected with *S. suis* type 2 (62, 64). To counter the low immunogenicity of CPSs, researchers have developed a conjugate vaccine composed of CPS linked to a carrier protein, as is the case for GBS. However, the efficacy of in-trial GBS conjugate vaccines depends on the CPS serotype included in the preparation. Indeed, CPSs of types Ia, Ib, II, and III induce strong protective IgG responses, whereas GBS type V CPS promotes higher concentrations of specific IgM than IgG (65). This suggests that the composition and/or structure of CPSs could influence its immunogenicity. Certain structural features of CPSs, such as variations in repeating unit composition or glycosidic linkage positions, are susceptible to producing different immune responses (67–69).

Despite these observations, very few studies have been dedicated to the characterization of CPS activity on APCs and to the corresponding signaling mechanisms. Some bacterial CPS motifs, including sialic acid, are very similar to molecules expressed by human cells or tissues, which could result in immune evasion through molecular mimicry (69). Therefore, chemical alteration of sialic acid of *N. meningitidis* type B CPS by de-*N*-acetylation or deletion of this sugar in GBS type V CPS improved the immunogenicity of these two CPSs (50, 70). However, the consequences of capsular sialic acid manipulation on the capacity of CPS to acti-



**FIG 5** CCL2 production by DCs in response to stimulation by *S. suis* type 2 CPS is dose and time dependent. Native *S. suis* type 2 CPS (200, 100, 50, or 5  $\mu\text{g/ml}$ ) was incubated with DCs ( $10^6$  cells/ml). After 6 or 24 h, supernatants were collected and CCL2 levels were quantified by ELISA. Cells stimulated with medium alone served as a negative control ( $C^-$ ). Data are expressed as means with SEM (pg/ml) for at least three experiments with at least three technical replicates.



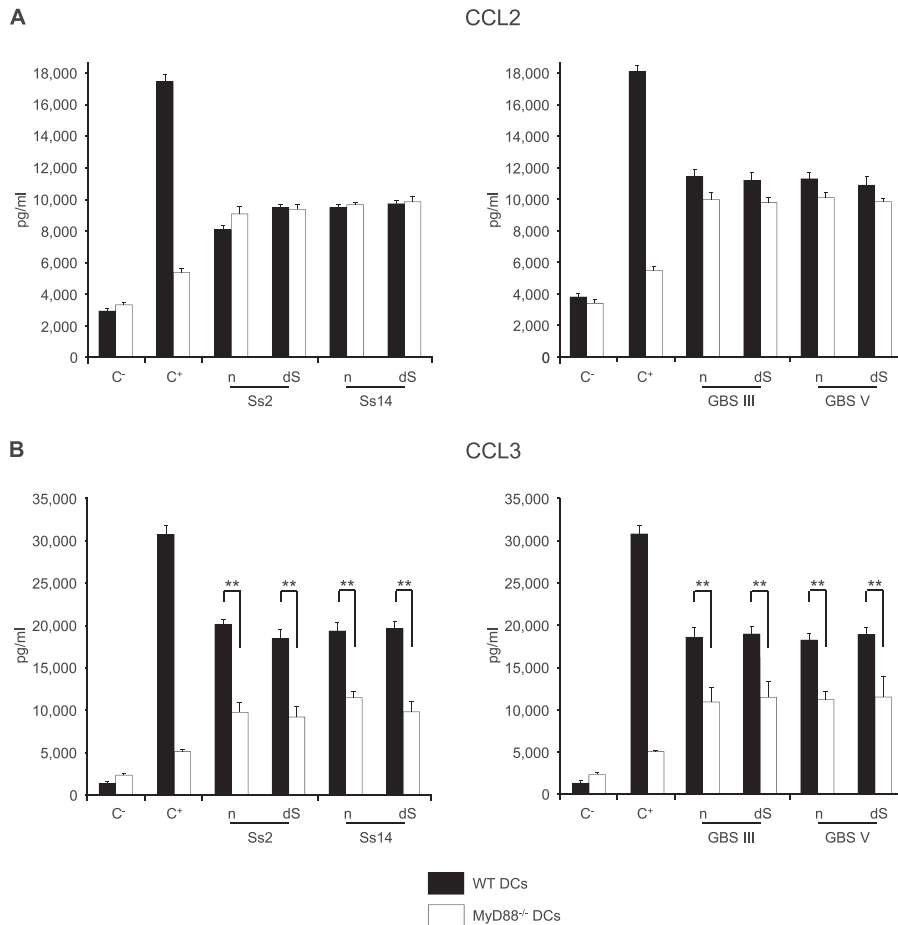
**FIG 6** Role of TLR2 in chemokine production by DCs in response to stimulation by *S. suis* or GBS CPSs for 24 h. Native (n) or desialylated (dS) *S. suis* type 2 (Ss2), *S. suis* type 14 (Ss14), GBS type III, or GBS type V CPSs (each at 200  $\mu$ g/ml) was incubated with WT (black bars) or TLR2<sup>-/-</sup> (white bars) DCs (10<sup>6</sup> cells/ml). After 24 h, supernatants were collected and CCL2 (A) and CCL3 (B) levels were quantified by ELISA. Cells stimulated with medium alone and with LPS (1  $\mu$ g/ml) served as negative (C<sup>-</sup>) and positive (C<sup>+</sup>) controls, respectively. Data are expressed as means with SEM (pg/ml) for at least three experiments with at least three technical replicates. \*,  $P < 0.05$ .

vate the immune system have been evaluated only in the context of Ab production. The mechanisms underlying this effect, such as the modulation of the function of APCs, are poorly known.

In this article, we analyzed the ability of *S. suis* and GBS CPSs to activate DCs as determined by production of cytokines and chemokines susceptible to being involved in TI responses. Mouse-origin DCs were used because mouse models are well described for both pathogens and because *in vitro* interactions of GBS or *S. suis* with mouse-, human-, and/or swine-origin cells show similar results (12, 14, 15, 53, 54, 71, 72). We used native and desialylated CPSs isolated from two of the most virulent and frequently isolated serotypes in humans for each bacterial species in order to examine the role of CPS composition in modulation of DC activation. The capsular preparations underwent a series of rigorous physicochemical and immunologic quality control tests, which attested the high purity of the CPSs, the preservation of epitope recognition, and the absence of sialic acid in the desialylated preparations. Our results demonstrated that native *S. suis* type 2 and 14 and GBS type III and V CPSs do not induce the release of key proinflammatory cytokines, which confirms the poor immunogenic nature of these molecules. This is in accordance with previous experiments where *S. suis* type 2 CPS did not induce the pro-

duction of either IL-1 $\beta$ , IL-6, or TNF- $\alpha$  by human monocytes or murine macrophages (53, 73). However, our results are in contradiction with earlier studies which showed that GBS type III CPS stimulated the production of TNF- $\alpha$  and IL-6 by human cord blood monocytes (74, 75). This difference could be explained by the use of different cells or by variations in the purification method. Nevertheless, the authors of those studies indicated that GBS type III CPS was a poor stimulator compared to other bacterial cell wall components (74, 75). Other studies using CPSs purified from other bacterial species have concluded likewise, that such a molecule is a poor activator of the immune system. *S. pneumoniae* CPS incubated with human monocyte-derived DCs produced no or very small amounts of IL-12 or IL-10 (76). Similarly, *N. meningitidis* type C CPS was unable to induce production of IL-6 and TNF- $\alpha$  by human monocytes or macrophages (77). By using desialylated CPSs, we demonstrated for the first time that sialic acid does not seem to play an inhibitory role in the release of proinflammatory cytokines by DCs. The presence of sialic acid has been linked to production of the regulatory cytokine IL-10 (78, 79). However, sialic acid in either GBS CPSs or *S. suis* CPSs did not result in increased IL-10 production by DCs. Thus, an imbalance between pro- and anti-inflammatory cytokine profiles cannot ex-





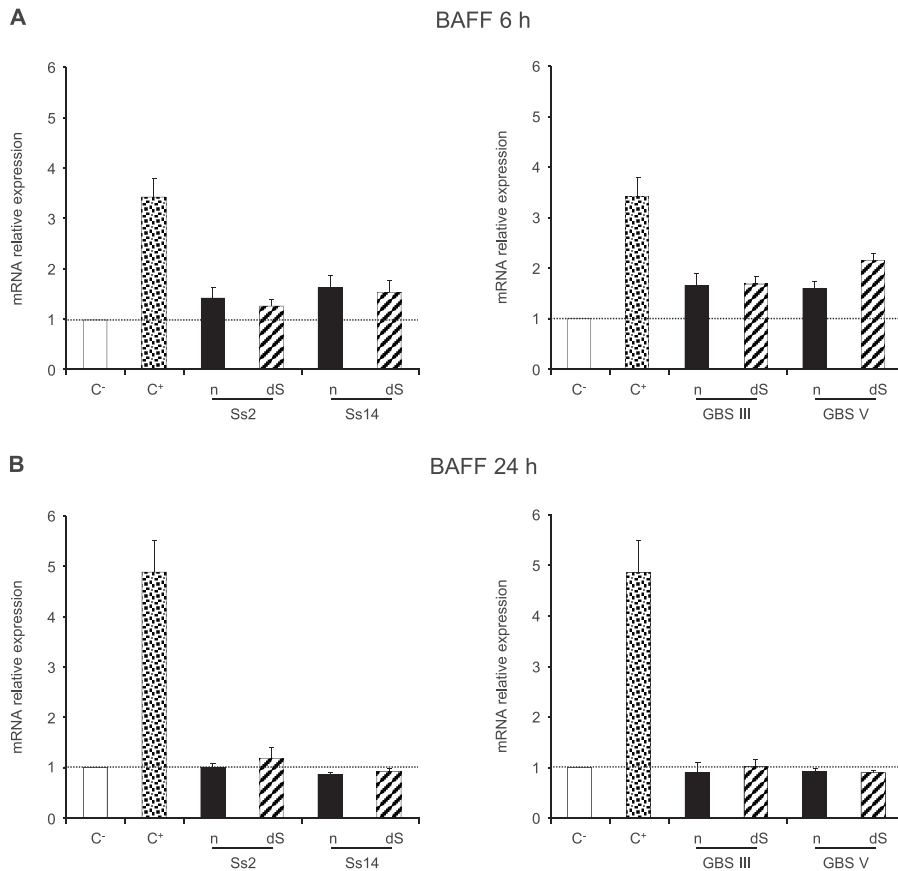
**FIG 7** Role of MyD88 in chemokine production by DCs in response to stimulation by *S. suis* or GBS CPSs for 24 h. Native (n) or desialylated (dS) *S. suis* type 2 (Ss2), *S. suis* type 14 (Ss14), GBS type III, or GBS type V CPS (each at 200  $\mu\text{g/ml}$ ) was incubated with WT (black bars) or MyD88<sup>-/-</sup> (white bars) DCs ( $10^6$  cells/ml). After 24 h, supernatants were collected and CCL2 (A) and CCL3 (B) levels were quantified by ELISA. Cells stimulated with medium alone and with LPS (1  $\mu\text{g/ml}$ ) served as negative (C<sup>-</sup>) and positive (C<sup>+</sup>) controls, respectively. Data are expressed as means with SEM (pg/ml) for at least three experiments with at least three technical replicates. \*\*,  $P < 0.001$ .

plain the poorly DC-stimulatory properties of either GBS or *S. suis* CPSs.

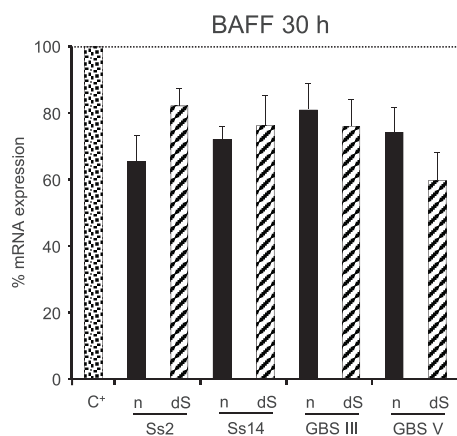
Interestingly, *S. suis* and GBS CPSs remarkably stimulated DC production of two members of the CC family of chemokines, i.e., CCL2 and CCL3. These two chemokines are known to play a major role in the selective recruitment of monocytes, macrophages, DCs, and lymphocytes to sites of inflammation (80, 81). A high level of CCL2 in the central nervous system is a characteristic of patients with bacterial meningitis (82). Systemic production of this chemokine, as well as its expression in the brain, is a feature of *S. suis* type 2-infected mice (83), and CCL2 has been associated with clinical signs of GBS sepsis in neonates (84). Recently, whole *S. suis* type 2 was shown to induce CCL2 production by murine DCs (12) and CCL3 production by total mouse splenocytes (unpublished data). Similarly, studies have shown that whole GBS type III stimulates CCL2 and CCL3 secretion by murine macrophages and/or DCs (85, 86). Our observations with purified CPSs allow a better interpretation of previous data showing different patterns of chemokine production obtained with total leukocytes, monocytes, and/or DCs cultured in the presence of whole *S. suis* type 2 or GBS types III and V and their respective nonencapsulated mutants. Indeed, in these studies, production of CCL2

and/or CCL3 was significantly diminished with bacteria lacking CPS (53, 72, 86), suggesting an important role of CPS in contributing to the production of these chemokines. Our observations are analogous to those of antecedent studies where *S. suis* type 2 CPS was able to induce the expression of CCL2 mRNA in a porcine whole-blood culture system and the release of this chemokine by human monocytes and murine macrophages (53, 72). Other studies have reported that purified CPSs from *N. meningitidis*, *Porphyromonas gingivalis*, and *Bacteroides fragilis* stimulate liberation of chemokines by murine macrophages or DCs (25–27). Surprisingly, differences in composition or structure between *S. suis* and GBS CPSs or the presence of sialic acid did not deeply influence the release of chemokine by DCs.

It is well known that PRRs, including TLRs, are involved in the activation of immune cells by encapsulated bacteria and/or their purified CPSs (27, 57, 87). Our results indicate that the production of CCL2 by native or desialylated *S. suis* or GBS CPSs is independent of TLR2- or MyD88-related pathways. Similarly, production of CCL2 by murine macrophages incubated with *S. suis* type 2 CPS has previously been shown to be TLR2 and MyD88 independent (53). Macrophage expression of CCL2 induced by whole GBS type III has also been reported to be TLR2 independent (55).



**FIG 8** Relative expression of BAFF mRNA by DCs in response to stimulation by *S. suis* or GBS CPSs for 24 h. Native (n) or desialylated (dS) *S. suis* type 2 (Ss2), *S. suis* type 14 (Ss14), GBS type III, or GBS type V CPS (each at 200  $\mu$ g/ml) was incubated with DCs ( $10^6$  cells/ml). After 6 h (A) or 24 h (B) of incubation, cells were collected and BAFF mRNA expression was determined by RT-qPCR. Cells stimulated with IFN- $\gamma$  (10 ng/ml) served as a positive control ( $C^+$ ). Data are expressed as means with SEM for at least three experiments and are relative to the level for DCs stimulated with medium alone ( $C^-$ ), which was arbitrarily fixed to 1.



**FIG 9** Effect of preincubation of DCs with *S. suis* or GBS CPSs on IFN- $\gamma$ -induced expression of BAFF mRNA. DCs ( $10^6$  cells/ml) were prestimulated with native (n) or desialylated (dS) *S. suis* type 2 (Ss2), *S. suis* type 14 (Ss14), GBS type III, or GBS type V CPS (each at 200  $\mu$ g/ml) for 6 h prior to incubation with IFN- $\gamma$  (10 ng/ml) for 24 h. Cells were then collected, and BAFF mRNA expression was determined by RT-qPCR. Cells prestimulated with medium alone before addition of IFN- $\gamma$  served as a positive control ( $C^+$ ). Data are expressed as means with SEM for at least three experiments and are relative to the  $C^+$  level, which was arbitrarily fixed to 100%.

While CCL2 and CCL3 are two members of the CC family of chemokines presenting similarities in the regulation of their synthesis as well as in their biological functions, CCL3 production induced by *S. suis* or GBS CPSs was significantly diminished with MyD88 $^{-/-}$  DCs and partially affected with TLR2 $^{-/-}$  DCs. Interestingly, and in contrast to the case for CCL2, macrophage expression of CCL3 induced by whole GBS type III has been reported to be TLR2 dependent (55). While differential expression of these two chemokines has been reported in other systems (88), the underlying regulatory mechanisms are unknown. Nevertheless, the partial inhibition of CCL3 production in TLR2 $^{-/-}$  or MyD88 $^{-/-}$  DCs suggests that other TLRs as well as MyD88-independent pathways may be implicated in chemokine release by *S. suis* or GBS CPSs. The concept of direct interaction of CPSs with TLRs is still controversial. Although we used highly purified CPS material undergoing strict quality controls, the possibility of undetected contamination by traces of lipoproteins cannot be excluded. Cells deficient in TLR2 might also fail to express adequate levels of a receptor that may be relevant for CPS recognition. Members of the large family of lectin receptors are other possible receptor candidates (78, 89) and warrant further investigations.

CPS antigens, with few exceptions, are considered TI antigens. It is well known that the TNF family member BAFF plays a crucial role in the immune response against these antigens. Whereas sev-

eral studies have shown that the TLR ligand LPS or the prototype TI antigen NP-Ficoll stimulates expression or production of BAFF by total splenocytes, macrophages, or DCs (29, 58, 60, 90), there is a relative paucity of evidence concerning BAFF induction by CPSs. In our study, we observed that *S. suis* and GBS CPSs were unable to induce a significant expression of BAFF by DCs in comparison with that in unstimulated cells. This is in accordance with a previous study where *N. meningitidis* type C and GBS type V CPSs did not promote the release of BAFF by murine DCs (58), and it confirms the poor immunogenicity of CPSs. We observed that CPSs presented a suppressive effect on the capacity of IFN- $\gamma$  to induce BAFF expression, and some CPSs were able to provoke an inhibition of up to 40%. Similarly, *N. meningitidis* type C and GBS type V CPSs have been shown to inhibit intracellular and extracellular levels of IFN- $\gamma$ -induced BAFF in murine DCs (58). We evaluated for the first time the influence of sialic acid on this effect. However, sialylation did not play a major role in modulation of BAFF expression. The negative regulatory function of CPSs is not observed only on BAFF mRNA synthesis. The presence of CPS impairs cytokine release by DCs and macrophages activated by *S. suis* (12, 15, 53, 71). The inhibitory effect of CPS is not specific to cytokine expression but affects other cell functions, such as the expression of costimulatory molecules and major histocompatibility complex class II by DCs (15). *S. suis* type 2 CPS has been shown to downmodulate phagocytosis by destabilizing lipid microdomains and inhibiting activation of signaling pathways involved in phagocytosis (91, 92), whereas GBS type III CPS impairs bactericidal functions of neutrophils (78).

In conclusion, we found that highly purified CPSs isolated from two distinct serotypes of two different Gram-positive streptococci, *S. suis* and GBS, were principally poorly immunogenic antigens. However, they were able to specifically induce production of CCL2 and CCL3 by DCs. TLR2 and other MyD88-dependent pathways are partially involved in recognition of these CPSs, which might also implicate a more complex cross talk with other receptors. Interestingly, the effect of CPS composition (including sialic acid) and structure on DC function was less marked than that previously reported for B cell activation (68, 69) or than the observed variations in CPS recognition by specific sera in our dot-ELISA analysis. Thus, DCs seem to recognize and respond to these CPSs in a “patterned” way rather than a structure-dictated manner, which is in agreement with the role of the innate immune system. Further studies on the impact of DC activation status on B cell responses to these TI antigens are guaranteed.

## ACKNOWLEDGMENTS

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